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Applicant : Fisher *et al.*

Serial No. : 09/724,028

Examiner: Tung, Joyce

Filed : November 28, 2000 Group Art Unit: 1637

For : METHOD FOR FULL-LENGTH cDNA CLONING USING  
DEGENERATE STEM LOOP ANNEALING PRIMERS

**BY HAND DELIVERY**

**February 26, 2003**

**LETTER**

Asst. Commissioner for Patents  
Box Missing Parts  
Washington DC 20231

Attention: Examiner Tung

Sir:

Enclosed herewith please find paper and computer-readable copies of a Revised Sequence Listing, and a Declaration requesting entry of the Revised Sequence Listing in the above-identified application. Entry of the Revised Sequence Listing should bring the above-identified application into full compliance with 37 C.F.R. §§1.821-1.825. Applicants apologize for the inadvertent omission of the four nucleotide sequences, now identified as SEQ ID NOS:30-33, from the response to the Notice to Comply with Sequence Requirements.

In light of the revisions to the Sequence Listing, please note that the primers depicted as D-SLAP, D-CLAP1, D-CLAP2 and T-SLAP in Figures 1B and C, to which Applicants intended to refer in the Response to Restriction Requirement filed on December 2, 2002, are now SEQ ID NOS:30-33, respectively. The species to be elected with traverse in this same Response to Restriction Requirement is now SEQ ID NO:30, rather than SEQ ID NO:1, as indicated in the Response.

Respectfully submitted,

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Enclosure